## **Amendments to the Specification:**

Please replace the paragraph beginning at page 1, line 3, with the following rewritten paragraph:

--The present This application is a continuation of U.S. Patent Application No. 09/843,676, filed April 26, 2001, pending, which is a continuation of U.S. Patent Application No. 08/854,050, filed May 9, 1997, now U.S. Patent No. 6,261,836; which is a continuation-in-part of U.S. Patent Application No. 08/851,843, filed May 6, 1997, now U.S. Patent 6,093,809; which is a continuation-in-part of U.S. Patent Application No. 08/846,017, filed April 25, 1997, now abandoned; which is a continuation-in-part of U.S. Patent Application No. 08/844,419, filed April 18, 1997, now abandoned; which is a continuation-in-part of U.S. Patent Application No. 08/724,643, filed October 1, 1996, now abandoned. Each of the aforementioned applications is explicitly incorporated herein by reference in its entirety and for all purposes.--

Please replace the paragraph beginning at page 9, line 15, with the following rewritten paragraph:

--In alternative preferred embodiments, the present invention provides polynucleotide sequences corresponding to the human telomerase, including SEQ ID NOS:173 and 224, and their complementary sequences. The invention further contemplates fragments of these polynucleotide sequence (*i.e.*, SEQ ID NOS: 173 and 224) that are at least 5 nucleotides, at least 20 nucleotides, at least 250 nucleotides, and at least 500 nucleotides in length. The invention further contemplates fragments of the complements of these polynucleotide sequences (*i.e.*, SEQ ID NOS: 173 and 224) that are at least 5 nucleotides, at least 20 nucleotides, at least 250 nucleotides, and at least 500 nucleotides in length. In addition, the invention features polynucleotide sequences that hybridize under stringent conditions to SEQ ID NOS: 173 and 224, and/or fragments, and/or the complementary sequences thereof. The present invention further contemplates a polynucleotide sequence comprising the complement of the nucleic acids of SEQ ID NOS: 173 and 224, or variants thereof. In a further embodiment, the polynucleotide sequence comprises a purified, synthetic

nucleotide sequence corresponding to a fragment of SEQ ID NOS: 173 and 224, having a length of about ten to thirty nucleotides. The present invention further provides plasmid pGRN121 (ATCC accession ##20916), and the lambda clone 25-1.1 (ATCC accession #\_\_\_\_\_\_\_accession # 209024).--

Please replace the paragraph beginning at page 11, line 29, with the following rewritten paragraph:

--The present invention also provides methods for detecting the presence of nucleotide sequences encoding at least a portion of human telomerase in a biological sample, comprising the steps of, providing: a biological sample suspected of containing nucleic acid corresponding to the nucleotide sequence of SEQ ID NO: 100, and/or SEQ ID NO: 173, and/or SEQ ID NO: 224; the nucleotide polynucleotide of SEQ ID NO: 100, and/or SEQ ID NO: 173, and/or SEQ ID NO: 224, or fragment(s) thereof, combining the biological sample with the nucleotide polynucleotide under conditions such that a hybridization complex is formed between the nucleic acid and the nucleotide polynucleotide; and detecting the hybridization complex.--

Please replace the paragraph beginning at page 13, line 17, with the following rewritten paragraph:

--Figure 9 shows Figures 9A and 9B show the DNA sequence of the gene encoding the 123 kDa telomerase protein subunit (SEQ ID NO:1).--

Please replace the paragraph beginning at page 13, line 23, with the following rewritten paragraph:

--Figure 12 shows <u>Figures 12A-12D show</u> the DNA sequence, as well as the amino acid sequences of all three open reading frames of the 43 kDa telomerase protein subunit (SEQ ID NOS:4-6).--

Please replace the paragraph beginning at page 13, line 25, with the following rewritten paragraph:

--Figure 13 shows Figures 13A and 13B show a sequence comparison between the 123 kDa telomerase protein subunit of *E. aediculatus* (SEQ ID NO:2) and the 80 kDa polypeptide subunit of *T. thermophila* (SEQ ID NO:52).--

Please replace the paragraph beginning at page 13, line 28, with the following rewritten paragraph:

--Figure 14 shows Figures 14A and 14B show a sequence comparison between the 123 kDa telomerase protein subunit of *E.aediculatus* (SEQ ID NO:2) and the 95 kDa telomerase polypeptide of *T. thermophila* (SEQ ID NO:54).--

Please replace the paragraph beginning at page 14, line 32, with the following rewritten paragraph:

--Figure 30 shows Figures 30A and 30B show the DNA sequence of *tezl* (SEQ ID NO:68).--

Please replace the paragraph beginning at page 15, line 2, with the following rewritten paragraph:

--Figure 33 is Figures 33A and 33B present a schematic summary of the tezl<sup>+</sup> sequencing experiments.--

Please replace the paragraph beginning at page 15, line 7, with the following rewritten paragraph:

--Figure 36-Figures 36A and 36B (SEQ ID NOS:58, 118, 121-130) shows show the alignment of the M2 PCR product with *E. aediculatus* p123, *S. cerevisiae*, and *Oxytricha* telomerase protein sequences.--

Please replace the paragraph beginning at page 15, line 18, with the following rewritten paragraph:

--Figure 42 Figures 42A-42J (SEQ ID NOS:2, 55 and 69) shows show the alignment of three telomerase sequences.--

Please replace the paragraph beginning at page 15, line 24, with the following rewritten paragraph:

--Figure 46 shows Figures 46A-46F show the DNA (SEQ ID NO:68) and amino acid (SEQ ID NO:69) sequence of *tez1*, with the coding regions indicated.--

Please replace the paragraph beginning at page 15, line 26, with the following rewritten paragraph:

--Figure 47 shows Figures 47A-47C show the DNA (SEQ ID NO:100) and amino acid (SEQ ID NO:101) of the ORF encoding an approximately 63 kDa telomerase protein or fragment thereof.--

Please replace the paragraph beginning at page 15, line 31, with the following rewritten paragraph:

--Figure 50 provides Figures 50A and 50B provide the results of preliminary nucleic acid sequencing analysis of human telomerase (SEQ ID NO:173).--

Please replace the paragraph beginning at page 15, line 33, with the following rewritten paragraph:

--Figure 51 provides Figures 51A-51I provide the preliminary nucleic acid (SEQ ID NO:173) and deduced ORF sequences (SEQ ID NOS:174-223) of human telomerase.--

Please replace the paragraph beginning at page 16, line 2, with the following rewritten paragraph:

--Figure 53 provides Figures 53A-53F provide the nucleic acid (SEQ ID NO:224) and deduced ORF sequence (SEQ ID NO:225) of human telomerase.--

Please replace the paragraph beginning at page 16, line 4, with the following rewritten paragraph:

--Figure 54 provides a restriction map of lambda clone 25-1.1 (ATCC accession # accession # 209024).--

Please replace the paragraph beginning at page 82, line 8, with the following rewritten paragraph:

--In addition, human cDNA libraries (inserted into lambda) were probed with the EcoRI-NotI fragment of the clone (#AA281296). One lambda clone, designated "lambda 25-1.1," (ATCC accession #\_\_\_\_\_\_ accession # 209024) was identified as containing complementary sequences. Figure 54 shows a restriction map of this lambda clone. The human cDNA insert from this clone was subcloned as an EcoRI restriction fragment into the EcoRI site of commercially available phagemid pBluescriptIISK+ (Stratagene), to create the plasmid "pGRN121," which was deposited with the ATCC (ATCC accession #209016). Preliminary results indicated that plasmid pGRN121 contains the entire open reading frame (ORF) sequence encoding the human telomerase protein.--

## **Amendments to the Drawings:**

The attached sheets of drawings include changes to the following drawings as indicated below:

Fig. 9 (2 sheets) are replaced with attached sheets Figs. 9A and 9B;

Fig. 12 (4 sheets) are replaced with attached sheets Figs. 12A-12D;

Fig. 13 (2 sheets) are replaced with attached sheets Figs. 13A and 13B;

Fig. 14 (2 sheets) are replaced with attached sheets Figs. 14A and 14B;

Fig. 30 (2 sheets) are replaced with attached sheets Figs. 30A and 30B;

Fig. 36 (2 sheets) are replaced with attached sheets Figs. 36A and 36B;

Fig. 42 (1sheet) is replaced with attached sheets Figs. 42A-42J;

Fig. 46 (5 sheets) are replaced with attached sheets Figs. 46A-46F;

Fig. 47 (4 sheets) are replaced with attached sheets Figs. 47A-47C;

Fig. 50 (2 sheets) are replaced with attached sheets Figs. 50A and 50B;

Fig. 51 (10 sheets) are replaced with attached sheets Figs.51A-51I;

Fig. 53 (8 sheets) are replaced with attached sheets Figs. 53A-53E.

Attachment: Replacement Sheets

Annotated Sheet Showing Changes for Fig. 53